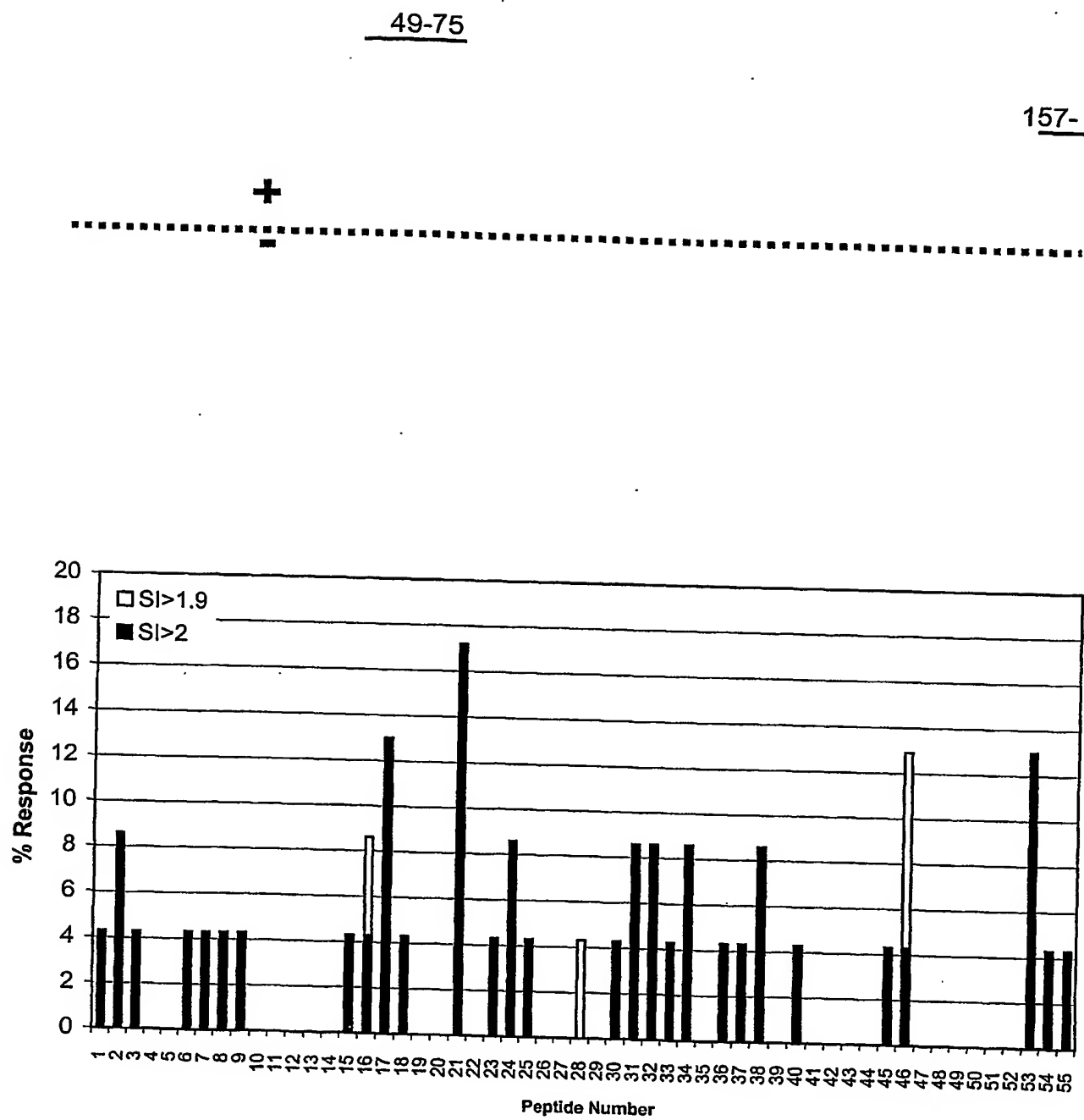


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Figure 1:**(A)****(B)**

SPAPPACDLRVL SKLLRDSHVLHS RLSQCPEVHPLP TPVLLPAVDFSL

1.

Region 1

GEWKTQMEETKA QDILGAVTLLLE GVMAARGQLGPT CLSSLLGQLSGQ

49

VRLLLGALQSLL GTQLPPQGRRTA HKDPNAIFLSFQ HLLRGKVRFLML

97

Region 2

VGGSTLCVRRAP PTTAVPSRTSLV LTINEL

145

174

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Figure 2A

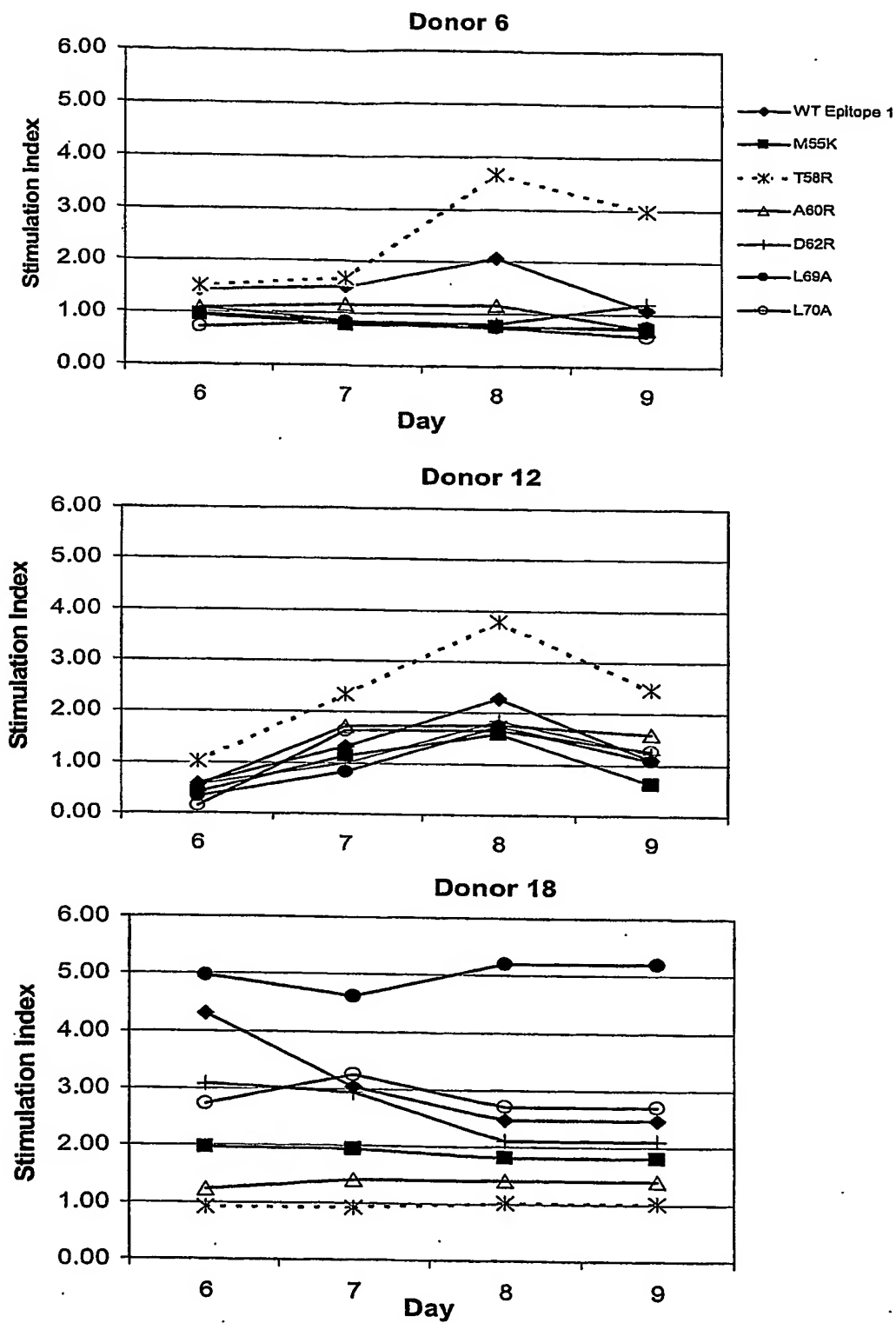


Figure 2B

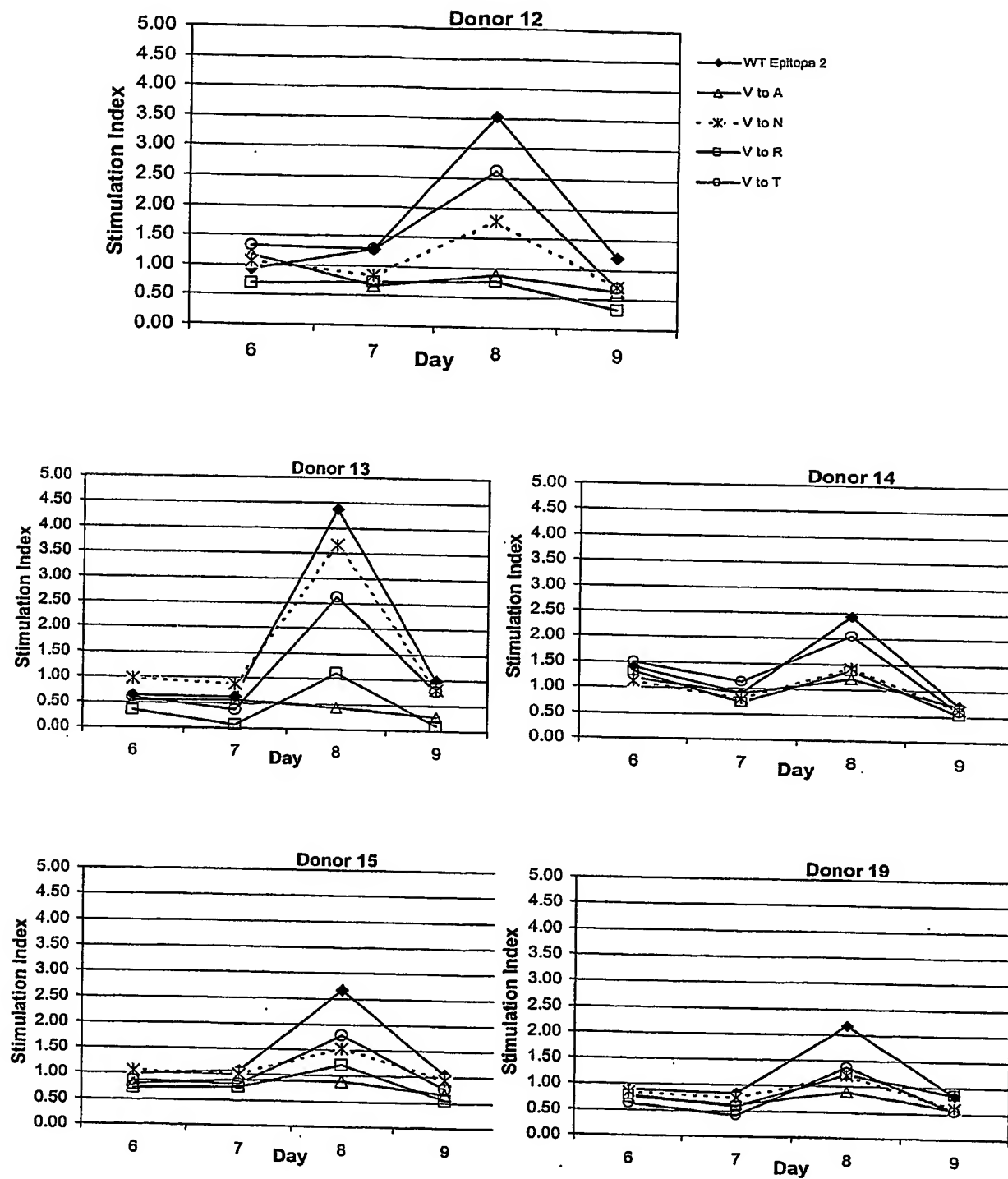


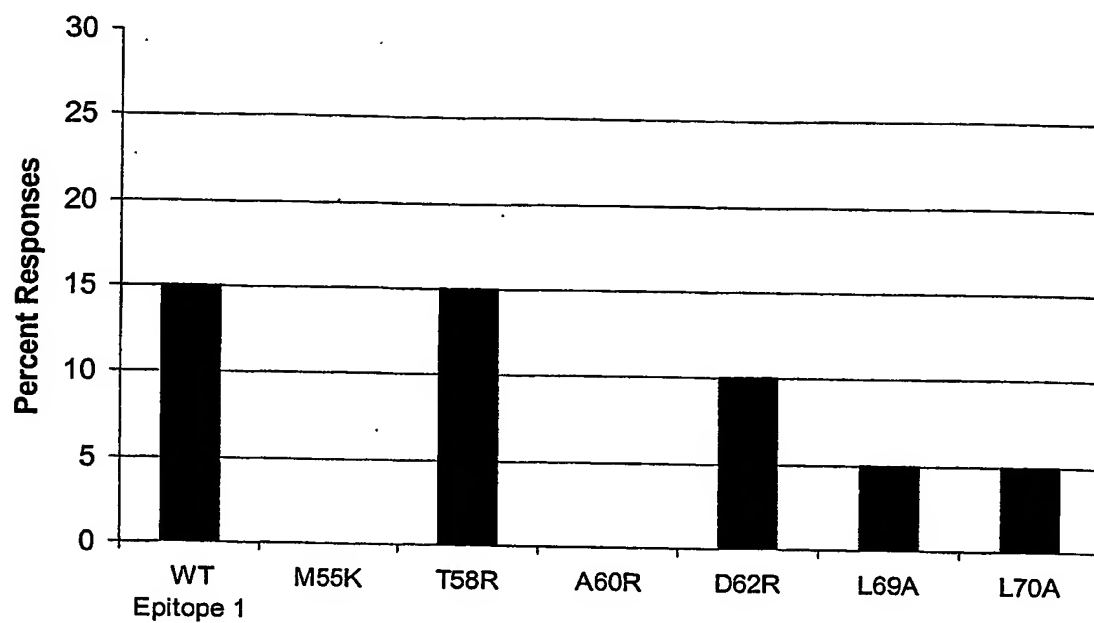
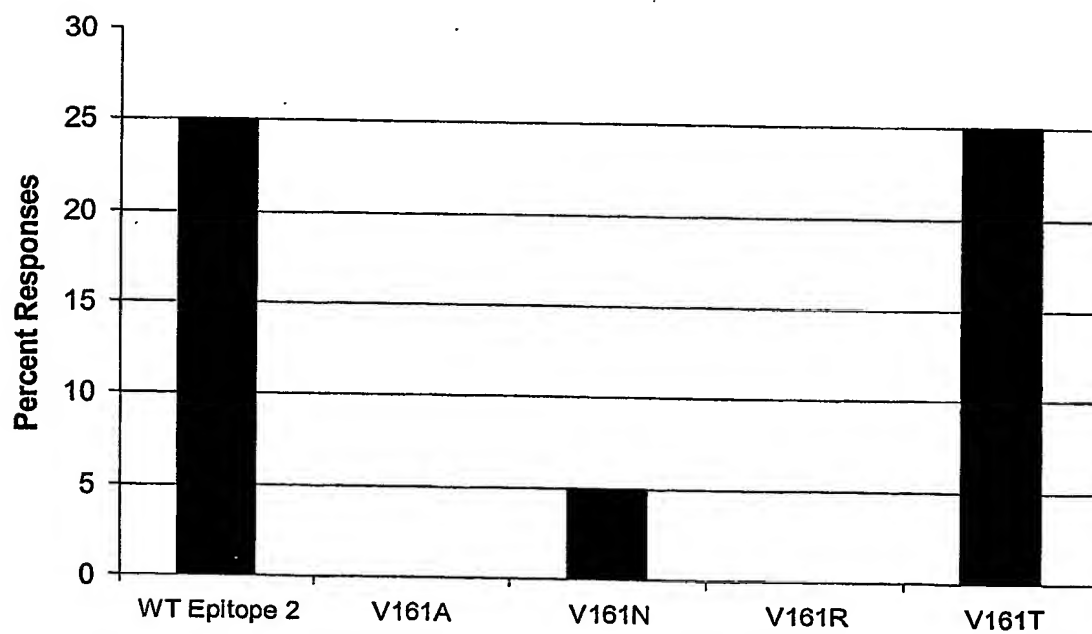
Figure 3**(A)****(B)**

Figure 4